VECTOR/PATHOGEN/HOST INTERACTION, TRANSMISSION

# Vector Competence of Selected African Mosquito (Diptera: Culicidae) Species for Rift Valley Fever Virus

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ABSTRACT Outbreaks of Rift Valley fever (RVF) in Egypt, Yemen, and Saudi Arabia have indicated the potential for this disease to spread from its enzootic areas in sub-Saharan Africa. Because little is known about the potential for most African mosquito species to transmit RVF virus (family Bunyaviridae, genus Phlebovirus, RVFV), we conducted studies to determine the vector competence of selected African species of mosquitoes for this virus. All eight species tested [Aedes palpalis (Newstead), Aedes mcintoshi Huang, Aedes circumluteolus (Theobald), Aedes calceatus Edwards, Aedes aegypti (L.), Culex antennatus (Becker), Culex pipiens (L.), and Culex quinquefasciatus Say], were susceptible to infection, and all except Ae. calceatus, Ae. aegypti and Cx. quinquefasciatus transmitted RVFV by bite after oral exposure. Estimated transmission rates for mosquitoes that successfully transmitted RVFV by bite ranged from 5% for Ae. mcintoshi to 39% for Ae. palpalis for mosquitoes that fed on a hamster with a viremia  $\ge 10^8$  plaque-forming units of virus/ml. We did not recover RVFV from any of 3,138 progeny of infected female mosquitoes. RVFV is unusual among arboviruses in that it has been isolated in nature from a large number of species and that numerous mosquitoes and other arthropods are able to transmit this virus in the laboratory. The recent introduction and spread of West Nile virus into the Americas and the spread of RVFV to the Arabian Peninsula illustrates the potential for viruses, once enzootic in Africa, to spread to other parts of the world.

KEY WORDS Rift Valley fever virus, transmission, mosquito, Africa, vector

Rift Valley fever virus (family *Bunyaviridae*, genus *Phlebovirus*, RVFV) has been associated with numerous outbreaks of severe disease in domestic ruminants in sub-Saharan Africa over the past 70 yr (Meegan and Bailey 1988, Gerdes 2004). However, the recent movement of RVFV out of Africa into the Arabian Peninsula (Jupp et al. 2002, Shoemaker et al. 2002, Balkhy and Memish 2003, Madani et al. 2003) has raised very real concerns regarding the agricultural and medical impact this zoonotic disease agent might have if it were to continue to spread (House et al. 1992). Although Rift Valley fever (RVF) is predominately a problem in domestic ruminants, where infections in pregnant an-

imals usually results in abortion and infection of newborn animals is nearly always fatal, humans are also susceptible to infection (Easterday et al. 1962, Meegan and Bailey 1988). In humans, most infections result in an undifferentiated febrile disease; however,  $\approx 1\%$  of the infections result in hemorrhagic complications, which are often fatal. In addition, ocular sequellae occur that can cause retinal damage, including blindness (Siam and Meegan 1980, Al-Hazmi et al. 2005).

Although RVFV is a member of the genus Phlebovirus and transmission by sand flies is known to occur in the laboratory (Hoch et al. 1984, Turell and Perkins 1990, Dohm et al. 2000), RVFV has been associated almost exclusively with mosquitoes in nature. It has been isolated from at least 40 species of mosquitoes in eight genera (Meegan and Bailey 1988, Fontenille et al. 1998). Laboratory studies have indicated that numerous species of mosquitoes are susceptible to oral infection and are able to transmit RVFV by bite (McIntosh et al. 1973b, 1980; Meegan and Bailey 1988, Gargan et al. 1988, Turell et al. 1996). However, some of these studies have focused on mosquitoes from areas where RVF is not enzootic in an attempt to determine the risk of local transmission of this virus, should it be introduced into a region where the mosquitoes are found (Gargan et al. 1988, Turell et al.

The opinions, interpretations, conclusions, and recommendations are those of the authors and are not necessarily endorsed by the U.S. Army.

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## **Report Documentation Page**

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14. ABSTRACT

Outbreaks of Rift Valley fever (RVF) in Egypt, Yemen, and Saudi Arabia have indicated the potential for this disease to spread from its enzootic areas in sub-Saharan Africa. Because little is known about the potential for most African mosquito species to transmit RVF virus (RVFV), we conducted studies to determine the vector competence of selected African species of mosquitoes for this virus. All eight species tested (Aedes palpalis (Newstead), Aedes mcintoshi Huang, Aedes circumluteolus (Theobald), Aedes calceatus Edwards, Aedes aegypti (L.), Culex antennatus (Becker), Culex pipiens (L.), and Culex quinquefasciatus Say, were susceptible to infection and all except Ae. calceatus, Ae. aegypti and Cx. quinquefasciatus transmitted RVFV by bite after oral exposure. Estimated transmission rates for mosquitoes that successfully transmitted RVFV by bite ranged from 5% for Ae. mcintoshi to 39% for Ae. palpalis for mosquitoes that fed on a hamster with a viremia >108 plaque-forming units of virus/ml. We did not recover RVFV from any of 3,138 progeny of infected female mosquitoes. RVFV is unusual among arboviruses in that it has been isolated in nature from a large number of species and that numerous mosquitoes and other arthropods are able to transmit this virus in the laboratory. The recent introduction and spread of West Nile virus into the Americas and the spread of RVFV to the Arabian Peninsula illustrates the potential for viruses, once enzootic in Africa, to spread to other parts of the world.

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Table 1. Source and colonization history of mosquitoes evaluated for their vector competence for Rift Valley fever virus

Species	Location	Yr	Generation
Ae. aegypti	Kenya	1982	P. "
Ae. calceatus	Kenya	1982	P."
Ae. circumluteolus	Kenya	1985, 1989	$\mathbf{P}_{0}^{0}/\mathbf{F}_{1}^{b}$
Ae. mcintoshi	Kenya	1985, 1986,	$\mathbf{P}_{a}/\mathbf{F}_{1}^{b}$
		1988	- 0 1
Ae. palpalis	Central African Republic	1985	$P_o/F_1^{\ b}$
Cx. antennatus	Kenya	1985	$\mathbf{p}^{-b}$
Cx. pipiens	Egypt	1980	Fb
Cx. quinquefasciatus	Kenya	1983	$rac{{{ m P}_o}^b}{{{ m F}_o}^{ ilde{b}}}$

<sup>&</sup>quot;Collected as eggs in Africa and reared to adults at USAMRIID.

1988a, Turell and Kay 1998). In addition, some of the studies with mosquitoes from areas where RVF is enzootic used large pools of mosquitoes (Smithburn et al. 1949; McIntosh et al. 1973b, 1980). Although these studies can determine whether a particular species is competent, they are unable to differentiate a highly efficient vector from a vector that is only marginally competent.

In our study, conducted during the 1980s, we examined eight species of mosquitoes collected in RVF enzootic areas for their susceptibility to oral infection and their subsequent ability to transmit RVFV by bite. Several of these species also were tested for their ability to vertically transmit RVFV to their progeny.

#### Materials and Methods

Mosquitoes. The mosquito species evaluated for their vector competence for RVFV and colonization histories are listed in Table 1. Mosquitoes were captured in Africa and transported to a biological safety level-3 laboratory (with HEPA-filtered exhaust air, treated sewage, and a 100% clothing change) at the United States Army Medical Research Institute of Infectious Diseases (USAMRIID). They were then provided apple slices as a carbohydrate source and held at 26°C for 7-10 d until either exposed to viremic hamsters or allowed to feed on an uninfected hamster to stimulate egg production. In addition to the fieldcollected female mosquitoes, first-generation progeny of some of these mosquitoes also were used in these studies. All larvae were reared under standard conditions at 26°C (Gargan et al. 1983).

In addition to the species tested for vector competence, Eretmapodites quinquevittatus Theobald, derived from specimens collected in South Africa, were tested for their ability to transmit RVFV vertically to their progeny

Viruses and Virus Assays. Three strains of RVFV: ZH501, isolated in 1977 from the blood of a 10-yr-old Egyptian girl who had a fatal RVFV infection (Meegan 1979); Zinga (DakArB1976), isolated from Mansonia africana (Theobald) mosquitoes captured in the Central African Republic in 1969; and a Kenyan strain (21445) isolated from Aedes mcintoshi Huang in 1983 were used throughout this study.

Individual specimens (mosquito larvae, pupae, or adults) were triturated in 1 ml of diluent (10% heatinactivated fetal bovine serum in Medium 199 (Invitrogen, Carlsbad, CA) with Hanks' salts and antibiotics) and frozen at -70°C until tested for infectious virus by a plaque-assay on Vero cell monolayers. Serial 10-fold dilutions of each specimen were tested on 12-well plates as described by Gargan et al. (1983). Virus titers were expressed as  $\log_{10}$  plaque-forming units (PFU) per specimen.

**Determination of Vector Competence.** To provide a source of viremic blood, adult female Syrian hamsters were inoculated intraperitoneally with 0.2 ml of a suspension containing  $\approx 10^4$  PFU of RVFV. These hamsters were anesthetized 1 or 2 d later and placed individually (i.e., one per cage) on the top of cages containing 50-150 mosquitoes. Immediately after mosquito feeding, 0.2 ml of blood was obtained from each hamster by cardiac puncture, and it was added to 1.8 ml of diluent. The blood suspensions were frozen at -70°C until assayed on Vero cell monolayers to determine the viremias at the time of mosquito feeding. In addition to the blood sample, three mosquitoes from each replicate were triturated individually in 1 ml of mosquito diluent immediately after feeding. These suspensions were tested by plaque assay to determine the actual virus dose ingested. After exposure to the viremic hamsters, engorged mosquitoes were transferred to 3.8-liter screen-topped cardboard cages. Apple slices, or a 7% sucrose solution, were provided as a carbohydrate source, and mosquitoes were held at 26°C and a photoperiod of 16:8 (L:D) h until tested for infection, dissemination, and transmission rates. Approximately 1 wk after the infectious bloodmeal, moist toweling or a water dish was added to each cage to stimulate oviposition.

To determine whether the mosquitoes could transmit virus by bite, mosquitoes were allowed to feed on susceptible hamsters either individually or in small groups of two to five mosquitoes each. Because RVFV infection consistently is fatal to hamsters, we considered death or euthanasia (when moribund) of these animals to indicate virus transmission. Presence of virus was verified by isolating virus from brain tissue from a subset of the dead hamsters. Immediately after each transmission trial, mosquitoes were killed by freezing at -20°C for 5 min, identified to species, and their legs and bodies triturated separately in 1 ml of diluent. These suspensions then were frozen at  $-70^{\circ}\mathrm{C}$ until tested for virus.

Mosquito infection was determined by recovering virus from its body tissue suspension. If virus was recovered from its body, but not its legs, the mosquito was considered to have a nondisseminated infection limited to its midgut. In contrast, if virus was recovered from both body and leg suspensions, the mosquito was considered to have a disseminated infection (Turell et al. 1984). The dissemination rate was the percentage of orally exposed mosquitoes that contained virus in their legs. Because some of the mosquitoes were tested for transmission in small pools, it was not always possible to determine which mosquito in a pool actually

<sup>&</sup>lt;sup>b</sup> Captured as adults.

Table 2. Infection and dissemination rates for mosquitoes orally exposed to Rift Valley fever virus

Species	• • •		······································			Day	s of ext	rinsic	incubati	on			
Species	Virus strain		3~10			11-16		≥18		3	Totals		
		N	$LR.^a$	D.R.b	N	$LR.^a$	$\mathbf{D}.\mathbf{R}.^{b}$	N	LR."	D.R.b	N	I.R.ª	D.R.
Infectious dose = $10^{5.8-6.8}$ PFU/ml					************		·						15.11.
Ae. circumluteolus	K-21445	10	70a	10a	19	58a	5) I	20					
Ae. mcintoshi	K-21445	83	35b	13a	57	23b	21ab	29	66a	34a	58	64a	26a
Ae. palpal <b>i</b> s	Zinga	NT	NT	NT	19	58a	7e	70	23c	10b	210	28Ь	10Ь
Cx. antennatus	K-21445	NT	NT	NT	17	ээа 18b	42a	34	56ab	41a	53	57a	42a
Infectious dose = $10^{7.0-7.8} \text{ PFU/ml}$				141	17	100	6bc	23	35bc	0Ь	40	28b	3Ь
Ae. aegypti	ZH501	2	100ab	0bc	3	33b	0	h Im					
Ae. calceatus	ZH501	13	92a	15bc	20	ააი 65ab	0c	NT	NT	NT	5	60bcd	0Ь
Ae. circumluteolus	ZH501	10	20c	10bc	10	30b	30bc	11	91ab	55a	44	80ab	32ab
Ae. mcintoshi	K-21445	20	60b	45ab	10	зоь 30b	Oc ani	6	17c	17a	26	23d	8b
Cx. quinquefasciatus	ZH501	12	67ab	8c	39	зор 33b	20bc	30	60bc	43a	60	55c	40a
Cx. pipiens	ZH501	50	74ab	8c	20	ээв 85a	5e	NT	NT	NT	51	41cd	6b
Infectious dose = 10 <sup>≤80</sup> PFU/ml		-517	1140	OC.	20	ола	<b>4</b> 5ab	NT	NT	NT	70	77ab	19b
Ae. aegypti	ZH501	41	85a	39ab	4	75abc	77 T	B 7000					
Ae. calceatus	ZH501	13	100a	3ab	20	19авс 100a	75ab 50ab	NT	NT	NT	45	84b	42b
Ae. circumluteolus	K-21445	11	73ab	36ab	22	82ab		11	100a	36Ь	44	100a	43b
Ae. mcintoshi	K-21445	115	55bc	34b	126	54c	59ab	9	67abc	44abc	42	76bc	50b
Ae. palpalis	Zinga	23	87a	61a	80	83a	43b	114	42c	18bc	355	50de	35b
Cx. antennatus	K-21445	25	52bc	8cd	40	оза 63bc	70a	66	91a	77a	169	86b	72a
Cx. pipiens	ZH501	15	93a	33ab	34	91a	15c	70	62b	19c	135	60cd	16c
Cx. quinquefasciatus	ZH501	9	22c	11bc	13		29Ьс	15	87ab	47b	64	91ab	3 <b>4</b> b
			220	1100	10	31bc	0c	NT	NT	NT	22	27e	5c

NT, not tested; N, number tested.

transmitted virus by bite. Therefore, if more than one mosquito with a disseminated infection fed in a pool, data from that pool were not used to calculate the transmission rate, regardless of hamster survival.

Inoculated Mosquitoes. We also inoculated some of the mosquitoes (Rosen and Gubler 1974) to produce a cohort of mosquitoes with a known disseminated infection. These mosquitoes were then tested individually on susceptible hamsters to examine for the presence of a salivary gland barrier (Kramer et al. 1981, Turell and Bailey 1987).

Vertical Transmission. To test for the potential for vertical transmission, adult female mosquitoes of selected species were inoculated with RVFV, held for 7 d at 26°C, and then allowed to feed, en masse, on an anesthetized, naïve hamster. An oviposition dish was added 5 d later and eggs collected. Seven days after the first bloodmeal, the mosquitoes were provided a second naïve hamster, and eggs were collected as described above. In some cases, a third ovarian cycle of eggs was collected. Eggs from these mosquitoes with known disseminated infections were hatched and reared at 26°C. The progeny were tested either as pools of up to 25 fourth-stage larvae or pupae, or they were reared to the adult stage and then tested separately in pools of up to 25 males or females. All pools were triturated in 2 ml of diluent and then tested for RVFV by plaque assay.

This research was conducted in compliance with the Animal Welfare Act and other federal statutes and regulations in force at the time the work was done and adhered to the principles stated in the Guide for the Care and Use of Laboratory Animals, 1978 or 1985. The facility where this research was conducted is fully accredited by the Association for Assessment and Accreditation of Laboratory Animal Care International.

#### Results

Vector Competence. Viremias in the 39 hamsters used to expose mosquitoes to RVFV ranged from  $10^{5.8}$  to  $10^{10.0}$  PFU/ml of blood ( $10^{3.3}$ - $10^{7.5}$  PFU ingested per mosquito, respectively). Viremias induced by each of the three strains of RVFV were similar, with hamsters inoculated with the ZH501, K-21445, and Zinga strains of RVFV, each producing a mean viremia of  $10^{8.3}$  PFU/ml of blood. Because infection rates tended to increase with increasing virus dose ingested, we arbitrarily grouped the mosquitoes into those exposed to low ( $10^{5.8-6.8}$  PFU/ml), moderate ( $10^{7.0-7.8}$  PFU/ml), or high ( $\ge 10^8$  PFU/ml) viremias.

All eight species were susceptible to infection after ingesting RVFV, even at the lowest dose that a particular species was exposed (Table 2). Although all species became infected, different "barriers" were present in different species (Table 3). These ranged from a midgut infection barrier associated with low infection rates, midgut escape barrier in which only a small percentage of infected mosquitoes developed a disseminated infection, or a salivary gland barrier in which only a small percentage of those mosquitoes with a disseminated infection transmitted virus by bite when allowed to refed on a susceptible vertebrate host.

The two Ae. (Stegomyia) species tested, Aedes aegypti (L.) and Aedes calceatus Edwards, were both

<sup>&</sup>lt;sup>a</sup> Infection rate, percentage of mosquitoes containing virus in their bodies. Infection rates in the same virus dose group followed by the same by Discompanion of the property of the same by Discompanion of the property of the same by Discompanion of the property of the same by Discompanion of the same by D

<sup>&</sup>lt;sup>b</sup> Dissemination rate, percentage of mosquitoes containing virus in their legs. Dissemination rates in the same virus dose group followed by the same letter are not significantly different at  $\alpha \approx 0.05$ .

Table 3. Vector competence for mosquitoes fed on hamsters with RVFV viremias ≥10<sup>a.o</sup> PFU/ml

Species	No. tested	Infection rate"	Dissemination rate <sup>b</sup>	Dissemination (1) rate <sup>r</sup>	Transmission (D) rate <sup>d</sup>	Estimated transmission rate
Ae. aegypti	45	84	42	50 (38)	14 (7)	(
Ae. calceatus	44	100	43	43 (44)	0 (21)	
Ae. circumluteolus	42	76	50	66 (21)	21 (48)	<2
Ae. meintoshi	355	50	35	70 (177)	· · · · /	10
Ae. palpalis	159	86	72	82 (137)	14 (171)	5
Cx. antennatus	135	60	16	27 (81)	55 (33)	39
Cx. pipiens	64	91	34	, ,	84 (38)	13
Cx. quinquefasciatus	22	27	5	38 (58) 17 (6)	100 (28) NT	34 <5

NT, not tested.

highly susceptible to infection and virus dissemination. However, a salivary gland barrier existed in these species as only one of 28 of these mosquitoes with a disseminated infection transmitted virus when fed on a susceptible hamster (Table 3).

All three Ae. (Neomelaniconion) species tested, Aedes circumluteolus (Theobald), Ae. mcintoshi, and Ae. palpalis (Newstead), were moderately susceptible to infection and virus dissemination, with at least 50% of each species becoming infected after ingesting blood containing ≥108 PFU/ml of RVFV, and at least 50% of the infected mosquitoes developing a disseminated infection (Tables 2 and 3). Of these three species, Ae. palpalis consistently had higher infection and dissemination rates than the other two species at each of the virus doses tested. As with the Ae. (Stegomyia) spp. tested, there was evidence of a salivary gland barrier (Tables 3 and 4). However, these ranged from a major barrier with Ae. mcintoshi (only 14% of the mosquitoes with a disseminated infection transmitted virus by bite) to a more moderate one for Ae. palpalis, 55% transmitted) (Table 4).

All three Culex species tested, Culex antennatus (Becker), Culex pipiens (L.), and Culex quinquefas-

ciatus Say, were susceptible to RVFV. However, Cx, pipiens were significantly more susceptible to infection ( $\chi^2 > 19.3$ , df = 1, P < 0.001) and to virus dissemination ( $\chi^2 > 8.2$ , df = 1, P < 0.01) than were either of the other two Culex spp. tested. We were not able to determine whether Cx, quinquefasciatus could transmit virus by bite because none of the ones with a disseminated infection fed on a susceptible host. However, there was little evidence of a salivary gland barrier in either Cx, antennatus or Cx, pipiens, because 84 and 100%, respectively, of the refeeding mosquitoes with a disseminated infection of these two species, transmitted virus by bite.

Virus Titer Recovered from Mosquitoes. For all species tested, the mean titers of virus recovered from specimens with a nondisseminated infection were between 10- and 1,000-fold lower than those recovered from specimens of the same species with a disseminated infection (i.e., with virus detected in their legs) (Table 5). For nearly all species tested, more virus was recovered from the legs of mosquitoes with a disseminated infection, almost always  $\approx 10^{4.3}$  PFU per leg sample, than was recovered from the entire body of those individuals with a nondisseminated infection.

Table 4. Transmission rates for mosquitoes with a disseminated infection with RVFV after either oral exposure or intrathoracic inoculation

	<u> </u>		Route	e of infection			
Species		Oral <sup>a</sup>		noculated	Totals		
	N <sup>b</sup>	T.R. (N) <sup>c</sup>	$N^b$	T.R. (N) <sup>c</sup>	$N^b$	T.R. (N)	
Ae. aegypti	4	0(0)	3	33 (1)	-7	14 (1)	
Ae. calceatus	21	0 (0)	NT	NT NT	21	14 (1)	
Ae. circumluteolus	17	18 (3)	31	23 (7)		0 (0)	
Ae. mcintoshi	97	12 (12)	74		48	21 (10)	
Ae. palpalis	26	54 (14)	7.9	16 (12)	171	14 (24)	
Cx. antennatus		60 (3)	20	57 (4)	33	55 (18)	
Cx. pipiens			33	88 (29)	38	84 (32)	
		100 (8)	20	100 (20)	28	100 (28)	

T.R., transmission rate; NT, not tested.

<sup>&</sup>lt;sup>a</sup> Infection rate = percentage of mosquitoes containing virus in their bodies.

<sup>&</sup>lt;sup>b</sup> Dissemination rate = percentage of mosquitoes containing virus in their legs.

Dissemination (I) rate = percentage of infected mosquitoes containing virus in their legs (no. of infected mosquitoes) (i.e., lack of a midgut escape barrier).

<sup>&</sup>lt;sup>d</sup> Transmission (D) rate = percentage of refeeding mosquitoes with a disseminated infection that transmitted RVFV by bite (no. with a disseminated infection that fed) (i.e. lack of a salivary gland barrier) (from Table 5).

The estimated transmission rate for mosquitoes feeding on a viremia  $\geq 10^6$  PFU/ml = the percentage of mosquitoes which developed a disseminated infection with RVFV multiplied by the transmission rate for those individuals with a disseminated infection.

<sup>&</sup>lt;sup>a</sup> Mosquitoes with a disseminated infection (virus in their legs) after oral exposure to RVFV.

<sup>&</sup>quot;Number of mosquitoes that fed.

Percentage of mosquitoes that fed that transmitted virus (no. that transmitted virus).

Table 5. Viral titers in mosquitoes orally infected with RVFV (assayed ≥7 d after oral exposure)

Species	Nondisse	minated			-	Dissen	inated		
	No. tested	Body	No. tested	Body	Legs	No. tested	Transmitters	No. tested	Nontransmitter
Ae. aegypti	5	4.1 (0.3)**	5	5.6 (0.3)	4.3 (0.4)				. voneransmitter
Ae. calceatus	44	4.8 (0.5)	35	. ,	( )	0	NA	.3	5.6(0.4)
Ae. circumluteolus	41	/		5.6 (0.3)	4.3(1.2)	0	NA	21	5.7 (0.3)
		3.5 (1.0)	47	5.7(0.6)	4.3(0.7)	3	5.9 (0.3)	17	5.6 (0.6)
Ae. meintoshi	63	2.7(0.8)	144	5.6 (0.6)	4.3 (0.6)	6	5.8 (0.6)		( /
Ae. palpalis	34	3.9(0.7)	137	5.4 (0.4)	4.4 (0.5)	-	, ,	60	5.7 (0.5)
Cx. antennatus	60	2.7 (0.7)		,	. ( ,	14	5.6(0.2)	11	5.7(0.4)
		, ,	21	5.5(0.7)	4.0(0.9)	3	6.2(0.9)	3	5.0 (0.9)
Cx. pipiens	21	2.9 (0.7)	18	5.0 (0.3)	3.9 (1.0)	8	5.5 (0.2)	ő	NA

NA, not applicable.

For mosquitoes with a disseminated infection for each species tested, virus titers of mosquitoes transmitting virus by bite were not significantly different than those that failed to transmit virus by bite  $(t \le 1.6, df \ge 4, P >$ 0.21) (Table 5).

Vertical Transmission Studies. Despite testing >3,138 progeny of mosquitoes inoculated with RVFV, we did not detect evidence of vertical transmission in these specimens (Table 6). Testing a sample of the inoculated adult females indicated that all of them were infected with RVFV.

#### Discussion

All eight mosquito species tested in these studies were susceptible to infection with RVFV, and all except Ae. calceatus, Ae. aegypti, and Cx. quinquefasciatus transmitted RVFV by bite after oral exposure. Although all of the species were susceptible to infection, different "barriers" (i.e., midgut infection, midgut escape, and salivary gland; Kramer et al. 1981) seemed to be the determining factor of the vector competence for the various species. Cx. quinquefasciatus had a major midgut infection barrier as only 27% became infected, even at the highest viremia levels tested (≥108 PFU/ml). This is consistent with other studies that found that this species is a relatively poor vector of RVFV (Turell and Kay 1998; McIntosh et al. 1980; M.J.T., unpublished data). All of the other species tested were generally susceptible to oral infection, with infection rates ≥50% when they fed on a hamster with a viremia ≥108 PFU/ml. At this exposure dose, the Aedes species tested had only a moderate midgut escape barrier with virus disseminating to the hemocoel in 43– 82% of the infected specimens tested (Table 3). However, there was a more severe midgut escape barrier in the three Culex species, with only 17–38% of the infected

Table 6. Lack of vertical transmission of RVFV by mosquitoes

Species					
	Larval	Pupal	Male	Female	Totals
Ae. aegypti	495"	0	148	116	759
Ae. circumluteolus	0	0	55	102	157
Ae. mcintoshi	0	23	539	379	942
Er. quinquevittatus	728	0	334	218	1,280

<sup>&</sup>quot;Number of progeny (by stage) of RVFV-infected mosquitoes tested

specimens developing a disseminated infection. Therefore, a midgut escape barrier seemed to be the principal determinant of vector competency in the Culex species. This is similar to what has been reported for Cx. pipiens (Turell et al. 1984).

Our failure to demonstrate transmission of RVFV by Ae. aegypti and Cx. quinquefasciatus may have been due to the relatively small sample size tested as only four Ae. aegypti and no Cx. quinquefasciatus with a disseminated infection after oral exposure refed on a susceptible hamster. However, an inoculated Ae. aegypti mosquito in this study did transmit RVFV by bite, and orally exposed and inoculated Ae. aegypti and Cx. quinquefasciatus have been shown to be able to transmit RVFV (McIntosh et al. 1980, Turell and Bailey 1987, Turell and Kay 1998). Therefore, there does not seem to be an absolute salivary gland barrier in either of these species. However, previous studies (McIntosh et al. 1980, Turell and Bailey 1987, Gargan et al. 1988, Turell et al. 1988a), indicate that although Aedes (Stegomyia) spp. can become infected and develop a disseminated infection after oral exposure to RVFV, these species tend to be inefficient vectors due to a salivary gland barrier (Kramer et al. 1981). In our study, although the Aedes species tested were highly susceptible to infection and virus dissemination, these species were generally inefficient vectors due to a salivary gland barrier, with ≤21% of Ae. aegypti, Ae. calceatus, Ae. circumluteolus, and Ae. mcintoshi successfully transmitting RVFV by bite. However, virtually all of the Cx. antennatus and Cx. pipiens with a disseminated infection that fed on a susceptible hamster transmitted RVFV by bite. In addition to the two Culex species examined in the current study, other studies report essentially a lack of a salivary gland barrier in Culex zombaensis Theobald, Culex tarsalis Coquillett, and Culex annulirostris Skuse (Gargan et al. 1988; Turell and Kay 1998; M.J.T., unpublished data). Similarly, for most Aedes species, transmission rates for mosquitoes with a disseminated infection have generally been  $\leq$ 50%. In addition to the ones in the current study, these include Aedes albopictus (Say), Aedes canadensis (Theobald), Aedes triseriatus (Say), Aedes vexans (Meigen), Aedes sollicitans (Walker), Aedes taeniorhynchus (Wiedemann), Aedes fowleri (Charmoy), Aedes juppi McIntosh, Aedes caballus (Theobald), Aedes cantator (Coquil-

<sup>&</sup>quot;Mean (SD) of the log10 PFU per specimen.

lett), and Aedes excrucians (Walker) (McIntosh et al. 1980; Gargan et al. 1988; Jupp and Cornel 1988; Turell and Bailey 1987; Turell et al. 1988a, 1988b). Therefore, these studies suggest that a midgut escape barrier seems to be the principal determinant of vector competence in the Culex species, whereas a salivary gland barrier is the principal determinant in the Aedes species.

As mosquitoes were exposed to higher viral doses, not only were infection rates generally higher but also the percentage of infected individuals that developed a disseminated infection increased. Therefore, the midgut escape barrier seemed to be dose dependent, independent of the infection rate. Similar findings also have been reported for other mosquitoes exposed to RVFV (Turell et al. 1988b).

We examined the relationship between the amount of virus recovered from a mosquito and the various barriers. As expected, for each species, mosquitoes with a disseminated infection had significantly more virus than members of the same species without a disseminated infection. For most species, those with a disseminated infection contained at least 100- to 1.000fold more virus than their infected, but nondisseminated, cage mates. In contrast, we did not find a difference between the titers of mosquitoes with a disseminated infection that did or did not transmit virus by bite. Therefore, although total body titer was an excellent predictor of virus dissemination beyond the midgut, it had no predictive value to determine which mosquito with a disseminated infection would be able to transmit virus by bite.

For each species tested, the transmission rate for mosquitoes with a disseminated infection after oral exposure was not significantly different ( $\chi^2 \leq 2.54$ ,  $df = 1, P \ge 0.11$ ) from that in those with a disseminated infection after intrathoracic inoculation. This allowed us to use animals more efficiently to obtain data about a possible salivary gland barrier in these species because all of the inoculated specimens were known to have a disseminated infection, and feeding success was greater in those specimens that did not have to take an "infectious" bloodmeal before the transmission attempt. A similar lack of differences in the transmission rates for mosquitoes with a disseminated infection after oral exposure to RVFV compared with those inoculated with this virus also has been reported for Ae. albopictus, Ae. fowleri, Aedes caspius (Pallas), Anopheles pharoensis Theobald, and Culex perexiguus Theobald (Turell et al. 1988a, 1988b, 1996). This has allowed us to calculate an estimated transmission rate (i.e., percentage of mosquitoes with a disseminated infection that transmit virus by bite multiplied by the percentages of mosquitoes that develop a disseminated infection after oral exposure) that should be an accurate estimate of the vector competence of that particular mosquito species. Our estimated transmission rates for those mosquito species that successfully transmitted RVFV by bite and fed on a hamster with a viremia ≥108 PFU of virus/ml ranged from 39% for Ae. palpalis to 5% for Ae. mcintoshi. Our "high" dose in this study, ≥108 PFU/ml is consistent with viremias

determined for natural infections with RVFV, where viremias in lambs and calves were up to  $10^{10.2}$  and  $10^{9.2}$  mouse intracranial LD<sub>50</sub>, respectively (Easterday 1965, McIntosh et al. 1973a), and viremias in humans were up to  $10^{5.6}$  mouse intracranial LD<sub>50</sub> (Meegan 1979). Therefore, the results obtained in our study should apply to these mosquito species when exposed to RVFV-infected cattle or sheep in a natural outbreak of RVF.

We did not recover RVFV from any of 3,138 progeny of infected female mosquitoes. This is consistent with the results of several previous studies that also failed to find laboratory confirmation of vertical transmission of RVFV (McIntosh et al. 1980, Jupp and Cornel 1988, Turell et al. 1988b). However, isolating RVFV from both male and female Ae. mcintoshi [reported as Aedes lineatopennis (Ludlow)] reared from field-collected larvae (Linthicum et al. 1985) clearly demonstrates that vertical transmission of this virus can occur under natural conditions. Additional studies are needed to further evaluate the potential for various mosquito species to maintain this virus vertically. Various studies have isolated RVFV from a number of mosquito species (Meegan and Bailey 1988). These studies include detection of RVFV from Ae. mcintoshi (as Ae. lineatopennis) (McIntosh 1972, Linthicum et al. 1985), Ae. circumluteolus (Kokernot et al. 1957), Ae. palpalis (Meegan and Bailey 1988), Cx. antennatus (Lee 1979), and Cx. pipiens (Meegan et al. 1980).

The recent introduction and spread of West Nile virus into the Americas and the spread of RVFV to the Arabian Peninsula illustrates the potential for viruses, once enzootic in Africa, to spread to other parts of the world. Additional studies are needed to evaluate other potential vectors of RVFV and to determine the role of other factors (e.g., environmental temperature) on the transmission of this pathogen.

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